

## SEQUENCE LISTING

<110>	Irvin, Randall T. Hodges, Robert S.		
<120>	Pseudomonas Treatment Composi	tion and Method	
<130>	0113190-00064		
<140> <141>	US 09/865,159 2003-03-01		
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Asp Gly Ser Cys Pro Ala Asn Thr Ala Ala Thr Ala Gly Ile Glu Lys
        35
Asp Thr Asp Ile Asn Gly Lys Tyr Val Ala Lys Val Thr Thr Gly Gly
Thr Ala Ala Ala Ser Gly Gly Cys Thr Ile Val Ala Thr Met Lys Ala
Ser Asp Val Ala Thr Pro Leu Arg Gly Lys Thr Leu Thr Leu
Gly Asn Ala Asp Lys Gly Ser Tyr Thr Trp Ala Cys Thr Ser Asn Ala
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Asp Asn Lys Tyr Leu Pro Lys Thr Cys Gln Thr Ala Thr Thr Thr
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Gly Ile Ala Gly Ser Lys Ile Lys Ile Gly Thr Thr Ala Ser Thr Ala
Thr Glu Thr Tyr Ala Gly Val Glu Pro Asp Ala Asn Lys Leu Gly Val
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Ile Ala Val Ala Ile Glu Asp Ser Gly Ala Gly Asp Ile Thr Phe Thr
Phe Gln Thr Gly Thr Ser Ser Pro Lys Asn Ala Thr Lys Val Ile Thr
Leu Asn Arg Thr Ala Asp Gly Val Trp Ala Cys Lys Ser Thr Gln Asp
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Pro Met Phe Thr Pro Lys Gly Cys Asp Asn
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actcctaaag atacccagta tgacattggc ttcaccgagt ctactttgct agatggttct
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ggtaagagtc agatccaggt aacggacaat aaagatggca ccgttgagtt ggtcgctacc
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ttgggtaaat cttctggttc cgccatcaaa ggggctgtaa tcactgtttc gcgtaaaaat
                                                                    300
gacggagtct ggaactgcaa aatcaccaaa actcctacag cttggaagcc caactacgct
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Gly Lys Glu Ile Val Ser Ser Ala Thr Pro Lys Asp Thr Gln Tyr Asp
        35
                            40
Ile Gly Phe Thr Glu Ser Thr Leu Leu Asp Gly Ser Gly Lys Ser Gln
Ile Gln Val Thr Asp Asn Lys Asp Gly Thr Val Glu Leu Val Ala Thr
                                        75
                    70
Leu Gly Lys Ser Ser Gly Ser Ala Ile Lys Gly Ala Val Ile Thr Val
Ser Arg Lys Asn Asp Gly Val Trp Asn Cys Lys Ile Thr Lys Thr Pro
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Thr Ala Trp Lys Pro Asn Tyr Ala Pro Ala Asn Cys Pro Asn Ser
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gctggtcaat gcgatccggg tgcgacgggt tccagtttgt tgactggtgc ttctcagact
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tctcaaaccc tgccaaccaa taccggtgtt ccgcaggttc tggatcctct gactactcaa
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accactatca ttgcgacttt tggtaacggc gcatccgcag ctatttctgg ccagactctg
                                                                     300
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acctqqactc qtqatqttaa tggtggctgg agctgtgcta ctaccgtaga tgctaaattc
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cgtcctaatg gctgtactga c
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<213>
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Gly Arg Thr Ala Val Gly Thr Ala Ala Gly Gln Cys Asp Pro Gly Ala
        35
Thr Gly Ser Ser Leu Leu Thr Gly Ala Ser Gln Thr Ser Gln Thr Leu
Pro Thr Asn Thr Gly Val Pro Gln Val Leu Asp Pro Leu Thr Thr Gln
                                        75
Thr Thr Ile Ile Ala Thr Phe Gly Asn Gly Ala Ser Ala Ala Ile Ser
                                    90
                85
Gly Gln Thr Leu Thr Trp Thr Arg Asp Val Asn Gly Gly Trp Ser Cys
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Ala Thr Thr Val Asp Ala Lys Phe Arg Pro Asn Gly Cys Thr Asp
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atcgaagccc taaaggccga gatagaagca cttaaggcag agatcgaggc gctaaaagcg
                                                                     120
                                                                     180
gaaatagagg ctctgaaggc aggcggtgga ggagaattcg ctcgttcgga aggcgcatct
gctcttgctt cggtcaatcc gttgaagact accgttgaag aggcgctttc tcgtggttgq
                                                                     240
agcgtgaaga gcggtacagg tacagaggac gctactaaga aagaggttcc tctgggggtg
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geggeagatg ctaacaaact gggtactate geacteaaac eegateetge tgatggtact
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gcagatatca ctttgacttt cactatgggc ggtgcaggac cgaagaataa agggaaaatt
                                                                     420
attaccctga ctcgtactgc agctgatggt ctctggaagt gcaccagtga tcaggatgag
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Ala Glu Ile Glu Ala Leu Lys Ala Glu Ile Glu Ala Leu Lys Ala Gly
Gly Gly Glu Phe Ala Arg Ser Glu Gly Ala Ser Ala Leu Ala Ser
Val Asn Pro Leu Lys Thr Thr Val Glu Glu Ala Leu Ser Arg Gly Trp
                                        75
                    70
Ser Val Lys Ser Gly Thr Gly Thr Glu Asp Ala Thr Lys Lys Glu Val
Pro Leu Gly Val Ala Ala Asp Ala Asn Lys Leu Gly Thr Ile Ala Leu
                                105
Lys Pro Asp Pro Ala Asp Gly Thr Ala Asp Ile Thr Leu Thr Phe Thr
                            120
        115
Met Gly Gly Ala Gly Pro Lys Asn Lys Gly Lys Ile Ile Thr Leu Thr
                        135
Arg Thr Ala Ala Asp Gly Leu Trp Lys Cys Thr Ser Asp Gln Asp Glu
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Gln Phe Ile Pro Lys Gly Cys Ser Arg
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gaagtatcag cacttgagaa gggcggtgga ggagaattcg ctcgttcgga aggcgcatct
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gctcttgctt cggtcaatcc gttgaagact accgttgaag aggcgctttc tcgtggttgg
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agcgtgaaga gcggtacagg tacagaggac gctactaaga aagaggttcc tctgggggtg
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gcggcagatg ctaacaaact gggtactatc gcactcaaac ccgatcctgc tgatggtact
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gcagatatca ctttgacttt cactatgggc ggtgcaggac cgaagaataa agggaaaatt
                                                                     420
attaccctga ctcgtactgc agctgatggt ctctggaagt gcaccagtga tcaggatgag
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Lys Glu Val Ser Ala Leu Glu Lys Glu Val Ser Ala Leu Glu Lys Gly
        35
Gly Gly Glu Phe Ala Arg Ser Glu Gly Ala Ser Ala Leu Ala Ser
Val Asn Pro Leu Lys Thr Thr Val Glu Glu Ala Leu Ser Arg Gly Trp
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Ser Val Lys Ser Gly Thr Gly Thr Glu Asp Ala Thr Lys Lys Glu Val
                                    90
Pro Leu Gly Val Ala Ala Asp Ala Asn Lys Leu Gly Thr Ile Ala Leu
                                105
Lys Pro Asp Pro Ala Asp Gly Thr Ala Asp Ile Thr Leu Thr Phe Thr
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Met Gly Gly Ala Gly Pro Lys Asn Lys Gly Lys Ile Ile Thr Leu Thr
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Arg Thr Ala Ala Asp Gly Leu Trp Lys Cys Thr Ser Asp Gln Asp Glu
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Gln Phe Ile Pro Lys Gly Cys Ser Arg
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atcgaagccc taaaggccga gatagaagca cttaaggcag agatcgaggc gctaaaagcg
                                                                    120
gaaatagagg ctctgaaggc aggcggtgga ggagaattcg cacgcgctca gcttagcgaa
                                                                    180
cgcatgaccc tggccagtgg tctcaagacg aaagtgagcg atatcttctc tcaggatggg
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                                                                    300
tcctgcccgg ctaatactgc tgccacggca ggcatcgaga aagataccga catcaacggc
                                                                    360
aagtatgttg ccaaggtaac aactggtggc accgcagctg cgtctggtgg ttgcactatc
gttgctacta tgaaagcctc tgatgtggct actcctctga gggggaaaac tctgactttg
                                                                     420
actctaggaa atgctgacaa gggttcttac acttgggcct gtacttccaa cgcagataac
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Gly Gly Glu Phe Ala Arg Ala Gln Leu Ser Glu Arg Met Thr Leu
Ala Ser Gly Leu Lys Thr Lys Val Ser Asp Ile Phe Ser Gln Asp Gly
                                        75
Ser Cys Pro Ala Asn Thr Ala Ala Thr Ala Gly Ile Glu Lys Asp Thr
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Asp Ile Asn Gly Lys Tyr Val Ala Lys Val Thr Thr Gly Gly Thr Ala
                                105
Ala Ala Ser Gly Gly Cys Thr Ile Val Ala Thr Met Lys Ala Ser Asp
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Val Ala Thr Pro Lys Arg Gly Lys Thr Leu Thr Leu Thr Leu Gly Asn
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Ala Asp Lys Gly Ser Thr Thr Trp Ala Cys Thr Ser Asn Ala Asp Asn
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Lys Tyr Leu Pro Lys Thr Cys Gln Thr Ala Thr Thr Thr Pro
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qaaqtatcaq cacttgagaa gggcggtgga ggagaattcg cacgcgctca gcttagcgaa
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                                                                    240
                                                                    300
tectgeeegg etaataetge tgeeaeggea ggeategaga aagataeega eateaaegge
aaqtatqttq ccaaqqtaac aactgqtggc accgcagctg cgtctggtgg ttgcactatc
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                                                                    480
actctaggaa atgctgacaa gggttcttac acttgggcct gtacttccaa cgcagataac
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Gly Gly Glu Phe Ala Arg Ala Gln Leu Ser Glu Arg Met Thr Leu
Ala Ser Gly Leu Lys Thr Lys Val Ser Asp Ile Phe Ser Gln Asp Gly
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Ser Cys Pro Ala Asn Thr Ala Ala Thr Ala Gly Ile Glu Lys Asp Thr
                                    90
Asp Ile Asn Gly Lys Tyr Val Ala Lýs Val Thr Thr Gly Gly Thr Ala
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Ala Ala Ser Gly Gly Cys Thr Ile Val Ala Thr Met Lys Ala Ser Asp
                                                125
                            120
Val Ala Thr Pro Leu Arg Gly Lys Thr Leu Thr Leu Thr Leu Gly Asn
                        135
Ala Asp Lys Gly Ser Tyr Thr Trp Ala Cys Thr Ser Asn Ala Asp Asn
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Lys Tyr Leu Pro Lys Thr Cys Gln Thr Ala Thr Thr Thr Pro
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gcgctggcga cgatcaaccc gctgaagacc actgttgaag agtcgctgtc gcgtggaatt
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gtcgagccgg atgccaacaa gttgggtgta attgctgtag caatcgaaga tagtggtgcg
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                                                                     420
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atcactctga accgtactgc ggatggggtc tgggcttgta aatctaccca ggatccgatg
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Ala Glu Ile Glu Ala Leu Lys Ala Glu Ile Glu Ala Leu Lys Ala Gly
        35
Gly Gly Glu Phe Ala Arg Ser Glu Gly Ala Ser Ala Leu Ala Thr
Ile Asn Pro Leu Lys Thr Thr Val Glu Glu Ser Leu Ser Arg Gly Ile
                                        75
Ala Gly Ser Lys Ile Lys Ile Gly Thr Thr Ala Ser Thr Ala Thr Glu
Thr Tyr Ala Gly Val Glu Pro Asp Ala Asn Lys Leu Gly Val Ile Ala
                                105
Val Ala Ile Glu Asp Ser Gly Ala Gly Asp Ile Thr Phe Thr Phe Gln
                            120
Thr Gly Thr Ser Ser Pro Lys Asn Ala Thr Lys Val Ile Thr Leu Asn
                        135
Arg Thr Ala Asp Gly Val Trp Ala Cys Lys Ser Thr Gln Asp Pro Met
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                    150
Phe Thr Pro Lys Gly Cys Asp Asn
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<210>
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gcqctggcga cgatcaaccc gctgaagacc actgttgaag agtcgctgtc gcgtggaatt
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gctggtagca aaattaaaat tggtactact gcttctactg cgaccgaaac atatgccggc
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gtcgagccgg atgccaacaa gttgggtgta attgctgtag caatcgaaga tagtggtgcg
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qqtqatatta cctttacctt ccagactggt acctctagtc ccaagaatgc tactaaagtt
atcactctga accgtactgc ggatggggtc tgggcttgta aatctaccca ggatccgatg
                                                                     480
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<211> 168

<212> PRT

<213> Pseudomonas aeruginosa

<400> 22

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Lys Glu Val Ser Ala Leu Glu Lys Glu Val Ser Ala Leu Glu Lys Gly 35 40 45

Gly Gly Glu Phe Ala Arg Ser Glu Gly Ala Ser Ala Leu Ala Thr 50 55 60

Ile Asn Pro Leu Lys Thr Thr Val Glu Glu Ser Leu Ser Arg Gly Ile 65 70 75 80

Ala Gly Ser Lys Ile Lys Ile Gly Thr Thr Ala Ser Thr Ala Thr Glu 85 90 95

Thr Tyr Ala Gly Val Glu Pro Asp Ala Asn Lys Leu Gly Val Ile Ala 100 105 110

Val Ala Ile Glu Asp Ser Gly Ala Gly Asp Ile Thr Phe Thr Phe Gln 115 120 125

Thr Gly Thr Ser Ser Pro Lys Asn Ala Thr Lys Val Ile Thr Leu Asn 130 135 140

Arg Thr Ala Asp Gly Val Trp Ala Cys Lys Ser Thr Gln Asp Pro Met 145 150 155 160

Phe Thr Pro Lys Gly Cys Asp Asn 165